

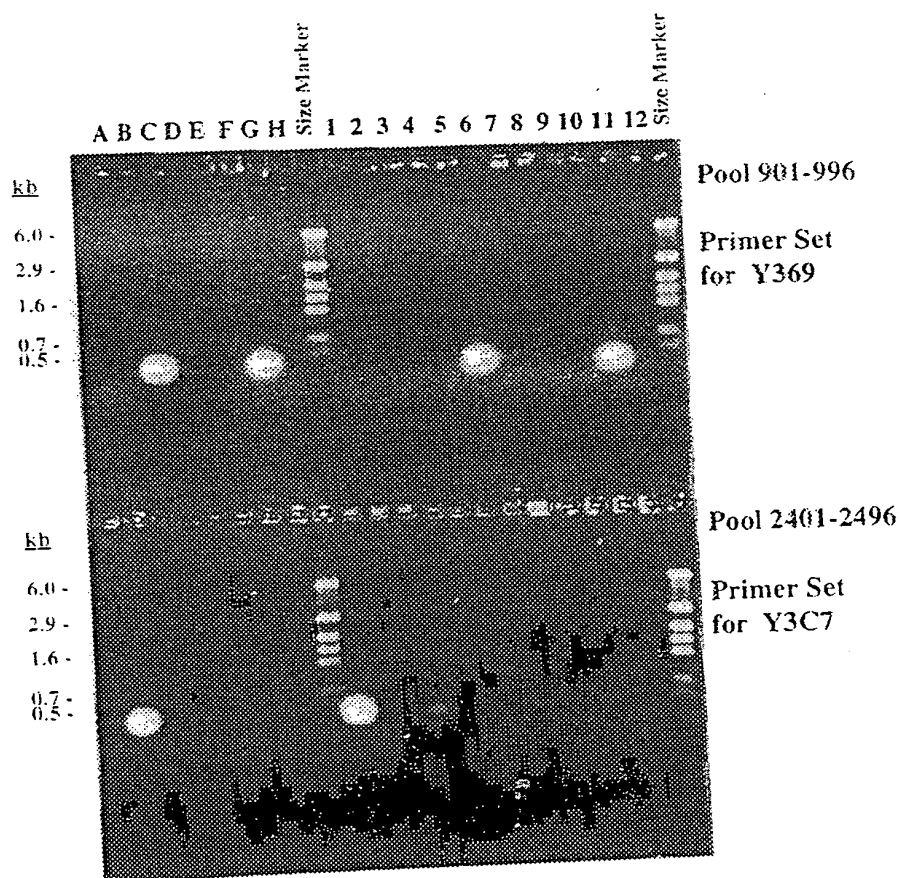
B

FIGURE 1B

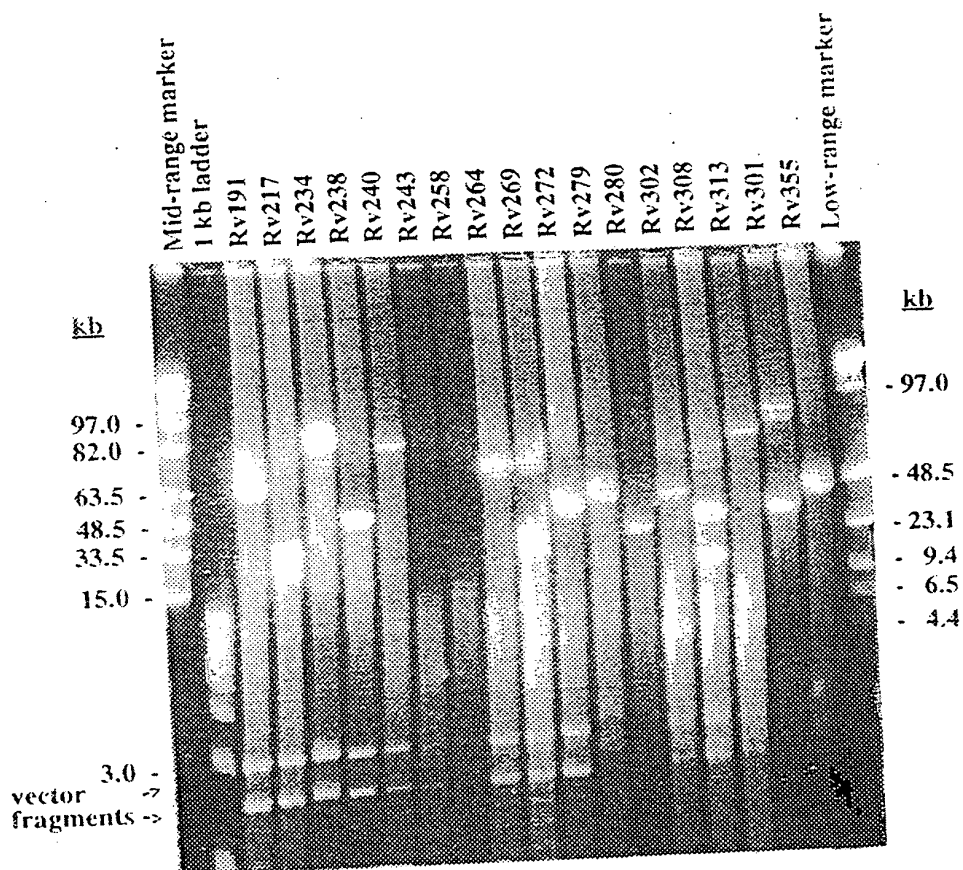


FIGURE 2

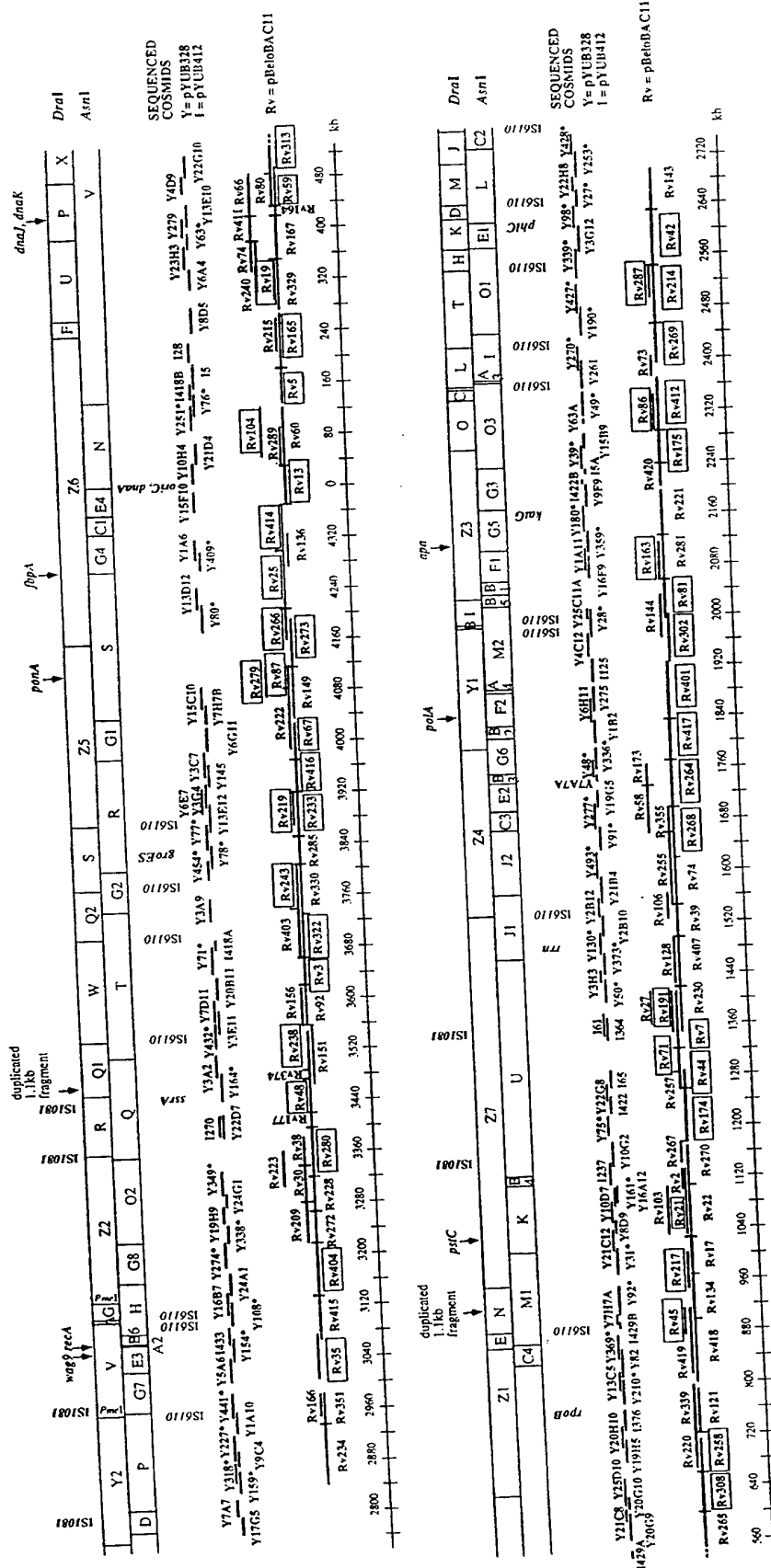


FIGURE 3

A

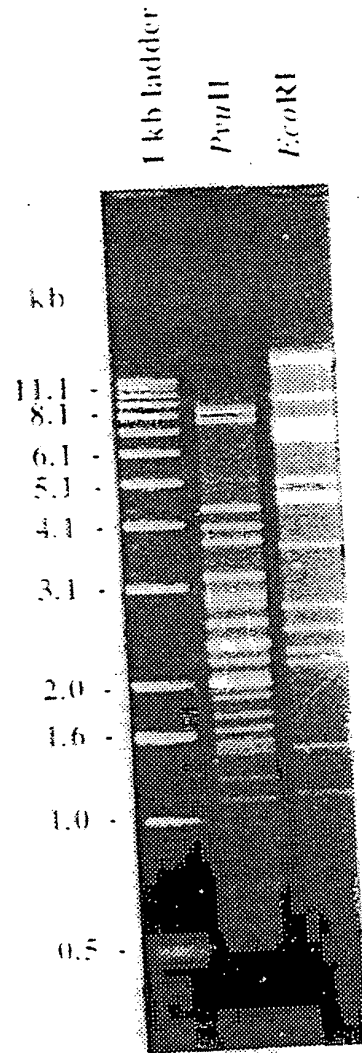


FIGURE 4A

B

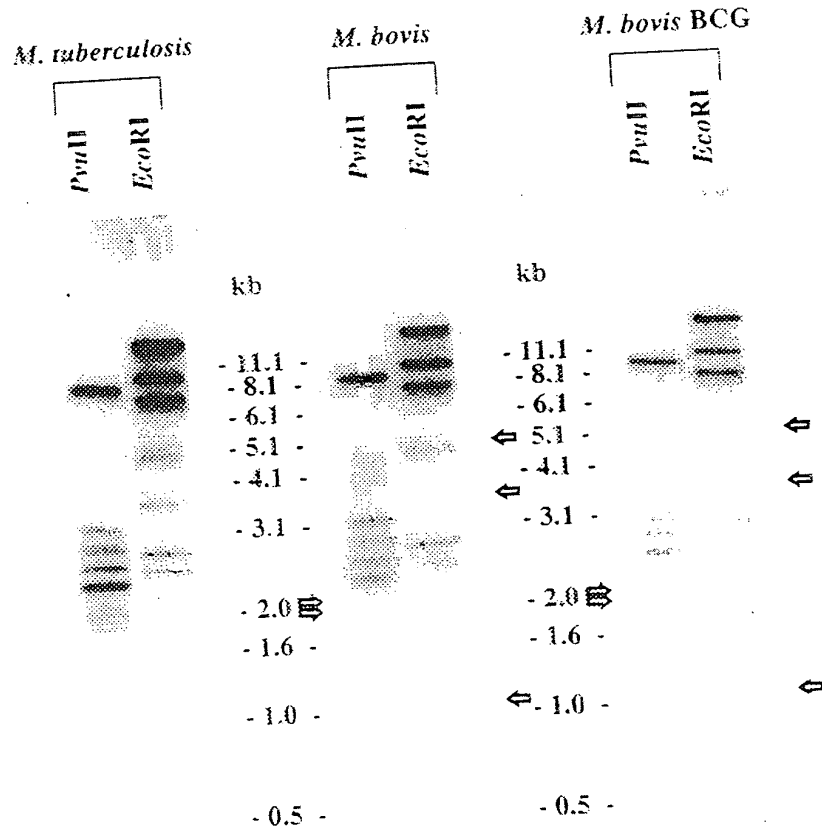


FIGURE 4B

[illegible]

Genomic region present in *M. tuberculosis* H37Rv only

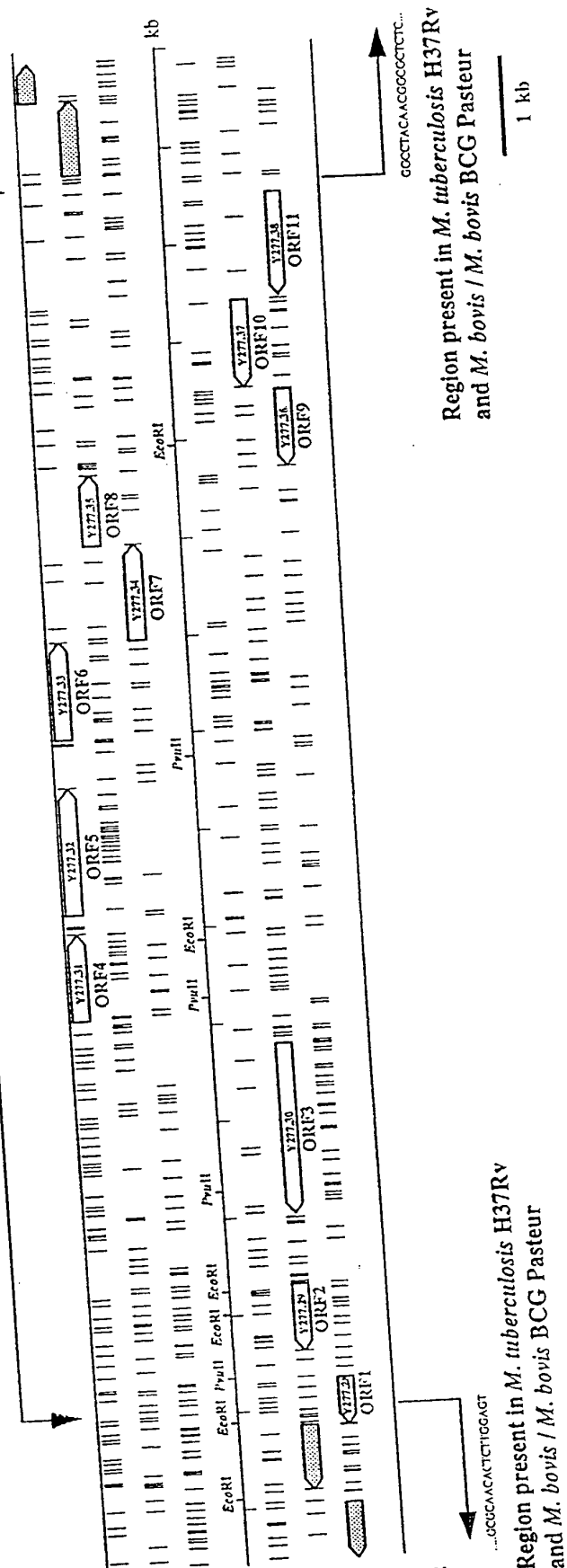


FIGURE 5

[illegible]

H37R+	PTQTLTGRPLIGNGTPGAYGSGATGAPGGWLLGDGGAGGSGAAAGSGAPGGAGGAAGLWGT	837273
BCG	PTQTLTGRPLIGNGTPGAYGSGATGAPGGWLLGDGGAGGSGAAAGSGAPGGAGGAAGLWGT	837453
H37R+	GGAGGAGGSSAGGGAGGAGGAGGWLLGDGGAGGIGGASTYLGTTGGGGYGGLWGAGGA	837633
BCG	GGAGGAGGSSAGGGAGGAGGAGGWLLGDGGAGGIGGASTYLGTTGGGGYGGLWGAGGA	
H37R+	GGAGGTGLYGGDGGAGGAGGTGGLLAGLIGAGGHHGTTGGLSTNGDGGYGGAGGNAGHLA	837813
BCG	GGAGGTGLYGGDGGAGGAGGTGGLLAGLIGAGGHHGTTGGLSTNGDGGYGGAGGNAGHLA	
H37R+	GPGGAGGAGGDGENLDTGGDGGAGGSAGLLFGSGGAGGAGGFGFLGGDGGAGGNAGLLLS	
BCG	GPGGAGGAGGDGENLDTGGDGGAGGSAGLLFGSGGAGGAGGFGFLGGDGGAGGNAGLLLS	
H37R+	SGGAGGFGGFTAGGYGGAGGNAGWLGF	837897
BCG	SGGAGGFGGFTAGGYGGAGGNAGWLGF	
H37R+	GGAGGYGGSAGLIGTGGNGGNGGTGANAGSPGTGGAGGLLLGQNGLNLGLP	838047
BCG	GGAGGYGGSAGLIGTGGNGGNGGTGANAGSPGTGGAGGLLLGQNGLNLGLP	
H37R+	GYEGGAALSYGDTGGAGGYGGSAGLIGTGGNGGNGGTGANAGSPGTGGAGGLLLGQNGLNLGLP	
BCG	GYEGGAALSYGDTGGAGGYGGSAGLIGTGGNGGNGGTGANAGSPGTGGAGGLLLGQNGLNLGLP	

FIGURE 6

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GCGGGCGGAA GGGGTTCGCG TCAGCGGGTG TTGGCGGGTG TCGGGGCTGG
NotI restriction site
 CTTAACTATG CGGCATCAGA GCAGATTGTA CTGAGAGTGC ACCATATGCG
 GTGTGAAATA CCGCACAGAT GCGTAAGGAG AAAATACCGC ATCAGGCGCC
 ATTCGCCATT CAGGCTGCGC AACTGTTGGG AAGGGCGATC GGTGCGGGCC
 TCTTCGCTAT TACGCCAGCT GCGGAAAGGG GGATGIGCTG CAAGGGGATT
 primer T7-BAC1
 AAGTTGGGTA ACGCCAGGGT TTTCCAGTC ACGACGTTGT AAAACGACGG
 CCAGTGAATTTGTAATACGAC TCAGTATAGG GCGAATTCGA GCTCGGTACC
 T7-promoter sequence
 CCGGGGATCCTCTAGAGTCGA CCTGCAGGCA TGCAGGCTTG AGTATTCAT
 primer T7-Belo2 *HindIII* cloning site SP6-promoter
 AGTGTCACCTAAATAGCTTG GCGTAATCAT GGTGATAGCT GTTCTGTG
 sequence (complementary strand) primer SP6-Mid (complementary strand)
 TGAAATTGTT ATCCGCTCAC AATCCACAC AACATACGAG CCGGAAGCAT
 AAAGTGTAAG GCCTGGGCTG CCAATGAGTGAGCTAACAC ACATTAATTG
 primer SP6-BAC1 (complementary strand)
 CGTTGCGCTC ACTGCCCCTG TTCCAGTCGG GAAACCTGTC GTGCCAGCTG
 CATTAATGAA TCGGCCAACG CGAACCCCTT GCGGGCGGCC GGGCCGTCGA
NotI restriction site

FIGURE 7